

Serial Number: 09/589,777A

ENTERED

BATCH 75

- ☐ Changed a file from non-ASCII to ASCII.
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically:
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____.
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____.
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____.
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____.
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____.
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____.
- ☐ Inserted mandatory headings, specifically: _____.
- ☐ Corrected an obvious error in the response, specifically: _____.
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____.
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____.
- ☒ Other: Seqs 2, 7, 13, 20 - inserted hard returns

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

1646

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/589,777A
 DATE: 12/26/2000
 TIME: 12:57:35

Input Set : A:\14401023011.txt
 Output Set: N:\CRF3\12262000\I589777A.raw

Does Not Comply
 Corrected Diskette Needed

4 <110> APPLICANT: Sukhatme, Vikas P.
 6 <120> TITLE OF INVENTION: Anti-Angiogenic Peptides and Methods of
 7 Use Thereof
 9 <130> FILE REFERENCE: 1440.1023-011
 11 <140> CURRENT APPLICATION NUMBER: US 09/589,777A
 12 <141> CURRENT FILING DATE: 2000-06-08
 14 <150> PRIOR APPLICATION NUMBER: PCT/US98/26057
 15 <151> PRIOR FILING DATE: 1998-11-16
 17 <150> PRIOR APPLICATION NUMBER: US 60/108,536
 18 <151> PRIOR FILING DATE: 1998-04-22
 20 <150> PRIOR APPLICATION NUMBER: US 60/082,663
 21 <151> PRIOR FILING DATE: 1998-04-22
 23 <150> PRIOR APPLICATION NUMBER: US 60/067,888
 24 <151> PRIOR FILING DATE: 1997-12-07
 26 <160> NUMBER OF SEQ ID NOS: 23
 28 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

97 <210> SEQ ID NO: 2
 98 <211> LENGTH: 184
 99 <212> TYPE: PRT
 100 <213> ORGANISM: Mus musculus
 102 <400> SEQUENCE: 2
 103 His Thr His Gln Arg Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
 104 1 10 15
 105 Thr Pro Leu Ser Glu Gly Met Arg Gly Ile Arg Gly Ala Asp Phe Gln
 106 20 25 30
 107 Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ser Gly Thr Phe Arg Ala
 108 35 40 45
 109 Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala
 110 50 55 60
 111 Asp Arg Gly Ser Val Pro Ile Val Asn Leu Lys Asp Glu Val Leu Ser
 112 65 70 75 80
 113 Pro Ser Trp Asp Ser Leu Phe Ser Gly Ser Gln Gly Gln Leu Gln Pro
 114 85 90 95
 115 Gly Ala Arg Ile Phe Ser Phe Asp Gly Arg Asp Val Leu Arg His Pro
 116 100 105 110
 117 Ala Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Ser Gly Arg
 118 115 120 125
 119 Arg Leu Met Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Thr Thr Gly
 120 130 135 140
 121 Ala Thr Gly Gln Ala Ser Ser Leu Leu Ser Gly Arg Leu Leu Glu Gln
 122 145 150 155 160
 E--> 123
 Lys Ala Ala Ser Cys His Asn Ser Tyr Ile Val Leu Cys Ile Glu Asn
 146 <210> SEQ ID NO: 5

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insert hard return

165

170

RAW SEQUENCE LISTING DATE: 12/26/2000
 PATENT APPLICATION: US/09/589,777A TIME: 12:57:35

Input Set : A:\14401023011.txt
 Output Set: N:\CRF3\12262000\I589777A.raw

147 <211> LENGTH: 24
 148 <212> TYPE: PRT
 149 <213> ORGANISM: Artificial Sequence
 151 <220> FEATURE:
 152 <223> OTHER INFORMATION: Leader peptide on protein produced by prokaryotic
 153 expression system pET17b, mouse endostatin begins
 154 immediately after.
 155 <400> SEQUENCE: 5

E--> 157

Met Gly His His His His His His His His His Ser Ser Gly His 1 5 10
 170 <210> SEQ ID NO: 7
 171 <211> LENGTH: 21
 172 <212> TYPE: PRT
 173 <213> ORGANISM: Artificial Sequence
 175 <220> FEATURE:
 176 <223> OTHER INFORMATION: Leader peptide on protein produced by prokaryotic
 177 expression system pET28a, mouse endostatin begins
 178 immediately after.
 180 <400> SEQUENCE: 7

E--> 181

Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro 1 5 10
 237 <210> SEQ ID NO: 13
 238 <211> LENGTH: 26
 239 <212> TYPE: PRT
 240 <213> ORGANISM: Artificial Sequence
 242 <220> FEATURE:
 243 <223> OTHER INFORMATION: Leader peptide on protein produced by eukaryotic
 244 yeast expression system pPIC2aA, mouse endostatin
 245 protein begins immediately after.
 247 <400> SEQUENCE: 13

E--> 248

Glu Phe Met Gly His His His His His His His His His Ser Ser 1 5 10
 315 <210> SEQ ID NO: 20
 316 <211> LENGTH: 8
 317 <212> TYPE: PRT
 318 <213> ORGANISM: Artificial Sequence
 320 <220> FEATURE:
 321 <223> OTHER INFORMATION: Leader peptide on protein produced by eukaryotic
 322 yeast expression system pPIC2aA, mouse endostatin
 323 protein begins immediately after.
 325 <400> SEQUENCE: 20

E--> 326 Glu Phe His His His His His His 1 5

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VERIFICATION SUMMARY
PATENT APPLICATION: US/09/589,777A DATE: 12/26/2000
TIME: 12:57:37

Input Set : A:\14401023011.txt
Output Set: N:\CRF3\12262000\I589777A.raw

L:123 M:252 E: No. of Seq. differs, <211>LENGTH:Input:184 Found:160 SEQ:2
L:157 M:252 E: No. of Seq. differs, <211>LENGTH:Input:24 Found:0 SEQ:5
L:181 M:252 E: No. of Seq. differs, <211>LENGTH:Input:21 Found:0 SEQ:7
L:248 M:252 E: No. of Seq. differs, <211>LENGTH:Input:26 Found:0 SEQ:13
L:326 M:252 E: No. of Seq. differs, <211>LENGTH:Input:8 Found:0 SEQ:20

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